Assessing species vulnerability to climate change

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Abstract

The effects of climate change on biodiversity are increasingly well documented, and many methods have been developed to assess species' vulnerability to climatic changes, both ongoing and projected in the coming decades. To minimize global biodiversity losses, conservationists need to identify those species that are likely to be most vulnerable to the impacts of climate change. In this review, we summarise different currencies used for assessing species' climate change vulnerability. We describe three main approaches used to derive these currencies (correlative, mechanistic and trait-based), and their associated data requirements, spatial and temporal scales of application and modelling methods. We identify strengths and weaknesses of the approaches and highlight the sources of uncertainty inherent in each approach that limit projection reliability. Finally, we provide guidance for conservation practitioners in selecting the most appropriate approach(es) for their planning needs and highlight priority areas for further assessments.

The Earth has warmed by about 0.74 °C in the last 100 years, and global mean temperatures are projected to increase further by 4.3 +/- 0.7 °C by 2100[1]. Agricultural expansion, overexploitation and invasive alien species introductions have been the main drivers of biodiversity loss in the recent past, but several lines of research suggest that climate change could become a prominent, if not leading cause of extinction over the coming century[2], both via direct impacts on species and through synergies with other extinction drivers[1,3]. Species have already responded to recent climatic shifts[4–8], and various attempts have been made to assess the potential risks to biodiversity posed by climate change over coming decades [9–11].

To assess the threats to a species posed by climate change one must have information regarding its vulnerability, which is defined by the IPCC as 'the predisposition to be adversely affected'[12]. Although there is currently no broad consensus in the scientific literature regarding the definition of 'species' vulnerability', it is generally accepted that this is a function of both intrinsic and extrinsic factors[13], and assessments often consider exposure, sensitivity and adaptability in combination[13,14]. Exposure is the magnitude of climatic variation in the areas occupied by the species[15]. Sensitivity, which is determined by traits that are intrinsic to species, is the ability to tolerate climatic variations, while adaptability is the inherent capacity of species to adjust to those changes[14,15]. Attempts at projecting the effects of climate change on species have used both different currencies (i.e. the range of measures used to assess species' climate change vulnerability) and divergent approaches for identifying the most vulnerable taxa. Because of this lack of consensus by the conservation community, a formal comparative evaluation is necessary to guide sensible choices of the most appropriate technique(s) for assessing species' vulnerability.

Here we provide the first comprehensive review of currencies and approaches that have been used to assess species' vulnerability to climate change, based on a total of 97 studies published between 1996 and 2014 (with >70% of the studies published during the last five years). We describe the four dominant currencies of species' climate change vulnerability assessments and

provide examples of how these have been applied. Three broad categories of approaches plus three combinations thereof were identified, and we describe each examining how they address uncertainties, and discuss their key limitations. Finally, we provide guidance for practitioners. Via these analyses, we aim to help conservationists select appropriate approaches for assessing species' vulnerability, such that climate change adaptation responses are as solidly based as possible.

Taxonomic and regional application of climate change vulnerability assessments of species

We conducted a systematic literature search using ISI Web of Knowledge. Key-words were selected to identify studies on climate change (climate change*, global warming*, sea-level rise*, elevated CO_2*, drought*, cyclones*, CO_2 concentration*) impacts (population reduction*, range changes*, range shift*, turnover*, extinction risk*, extinction probability*) that led to vulnerability assessments (vulnerability*, sensitivity*, adaptability*, exposure*) based on different types of approaches (mechanistic*, SDM*, correlative*, trait-based*, criteria*, niche models*). We then selected the most representative papers (in terms of both spatial and temporal scales, and taxa). Studies differed widely in taxonomic coverage, birds being the most frequently considered taxon, followed by mammals and plants, while non-insect invertebrates being seldom assessed (Fig. 1). Additionally, spatial scales of application and authors' interpretations of the concept of vulnerability varied extensively. More than 60% of the studies were developed at local scale, while only 4% of the papers assessed species' vulnerability globally (Fig. 1). As a result, numerous species have been assessed in only part of their range and their estimates of vulnerability may therefore be unrealistic.

Many published studies have shown that life-history traits are more important than taxonomy and distribution in determining species vulnerability to climate change[14]. Traits that commonly make a species vulnerable to climate change include limited dispersal abilities[14,16–18],

slow reproductive rates[11,19], specialised habitat and dietary requirements[14,20,21], restricted distribution and rarity[14,22], and narrow physiological tolerances[23–25], while potentially vulnerable habitats include intertidal areas, montane habitats, savannahs and grasslands[25]. Knowing what makes a species vulnerable and where vulnerable species are located can be very useful when practitioners need to assess the vulnerability of species for which only basic knowledge of their biology and ecology is available.

Studies conducted at a broad scale (regional, continental and global, almost 70% of the total), where used to derive a map of the areas with the greatest concentration of vulnerable species, according to an ecoregional classification (Fig. 2). For marine areas we performed a qualitative assessment (high, medium and low vulnerability, mostly based on Foden *et al.*[14]) because only a few marine taxa have been evaluated at broad scales and more than 80% of the species assessed are corals, while for terrestrial areas we were able to identify hotspots of vulnerable species as areas with high concentrations of vulnerable species (> 100), belonging to different taxonomic classes. These vulnerable areas, the Caribbean, the Amazon basin, Mesoamerica, eastern Europe through central and eastern Asia, the Mediterranean basin, the Himalayas, South-East Asia, North Africa, the Congo basin, tropical West Africa and Madagascar, should be a first priority for monitoring. However, over 70% of the studies we reviewed involved only three continents/subcontinents, with almost 33% of the studies in North America, 24% in Europe, and 14% in Australia (Fig. 3). By contrast, there is a paucity of studies in the most biodiverse tropical and subtropical regions of the world. Since climate change will act in concert with other threats, and habitat loss is predicted to severely affect biodiversity in developing countries[26], it is essential to conduct studies in these data deficient areas.

Currencies used to assess vulnerability: 'WHAT'

There is no standard way to assess a species' vulnerability to climate change, and the type of information (e.g. range extent, population size) needed will determine which approaches are most appropriate.

Distributional changes

To assess climate change impacts on species, current and future distributions can be projected using either mechanistic or correlative niche models (both approaches are discussed below), which relate environmental conditions to species' physiological responses or occurrence data, respectively. Several analyses have provided examples of species likely to suffer range reductions in the 21st century[16,18]. For example, Vieilleident *et al*.[27] predicted that the Malagasy baobab *Adansonia suarezensis* is likely to go extinct before 2080 due to an overall loss in suitable habitat. Changes in range size have usually been assessed by considering the climatic characteristics of current distributions and the projected distribution of these climatic conditions in future [27,28]. However, vulnerability might be exacerbated by other factors, including biotic interactions, reduced adaptive evolutionary response and dispersal ability. Several studies have incorporated dispersal ability into predictions of future range changes, either by contrasting scenarios of no dispersal with unlimited dispersal[29–31], by estimating average or maximum potential dispersal distances[16,18,24], or by explicitly simulating metapopulation dynamics including dispersal events[32,33]. For example, Schloss *et al.*[18] suggested that 87% of Western Hemisphere terrestrial mammals will likely experience a reduction in their climatically suitable area, with 20% of these species being particularly vulnerable due to their limited dispersal ability.

Population changes

A different set of modelling approaches uses predictions of population trends to inform risk assessments[34]. Quantified population changes can be based on direct observations, indices of abundance[34–36], reporting rates used as proxies for abundance[37], or they can be inferred from declines in extent of occupied or suitable habitat[34,38]. Examples of observed population declines within recent decades include long-distance avian migrants to Dutch forests, which have likely been driven principally by temperature changes in spring[35]. Also, a decrease in ice coverage has led to a reduction in polar bear (*Ursus maritimus*) numbers in the southern Beaufort Sea[39]. Some approaches to projecting future population sizes incorporate past population trends into mechanistic models[39–41], and consider the effects of changes in model parameters (e.g. distribution patterns, life history, climatic conditions). This type of approach has also been applied to a population of American marten (*Martes americana*) in North America, where explicit population models have been used to simulate a 40% decline in the population due to climate change by 2055[42].

Extinction probability

One synthesis estimated that between roughly 20 and 30% of species assessed are likely to be at increasingly high risk of extinction in the face of increasing global warming [12]. Extinction probability has been calculated for populations of species with known life-history characteristics, like the emperor penguin (*Aptenodytes forsteri*)[41], Arizona cliffrose (*Purshia subintegra*)[43], spring-summer chinook salmon (*Oncorhynchus tshawytscha*)[44] and polar bear (*Ursus maritimus*)[39], by using Population Viability Analyses[41,43], demographic models[39,44,45], or evolutionary models[46]. These methodologies combine population fluctuations with changing environmental parameters in order to estimate extinction probability within a given time interval. For example, Fordham *et al.*[45] modelled the predicted abundance of the Iberian lynx (*Lynx pardinus*) under three climate scenarios

193 by integrating temperature and precipitation data, prey availability and management interventions,
194 and predicted that climate change may drive this species to extinction within the next 50 years. This
195 work relied upon a thorough understanding of the species' biology and of demographic dynamics
196 related to extinction risk. However, as most species lack such detailed data, extinction risk due to
197 climate change tends to be quantified only for better-known species.

198

199 **Vulnerability indices and other relative scoring systems**

200 Vulnerability indices are quantitative indicators of the relative vulnerability of species. The
201 data derived from the currencies discussed above, and from trait-based vulnerability assessments
202 (TVAs), can be used to obtain scores[14], categories[34] or indices[47], which are often easier for scientists
203 and practitioners to interpret and use, in order to identify species at risk within their focal areas.
204 Foden *et al*.[14], for example, classified birds, amphibians and corals into two vulnerability categories
205 (low or high). One limitation of indices and scores is that they do not provide any direct measures
206 of the expected impact on species, i.e. they are not expressed in terms of any of the currencies
207 otherwise used to assess species' vulnerability (e.g. range reductions, extinction probability,
208 population decline).

209

210 **Approaches used to model species' vulnerability to climate change: 'HOW'**

211 Different approaches are used to assess species' vulnerability to climate change. These
212 approaches can be placed in four classes: 1) correlative, 2) mechanistic, 3) trait-based, and 4)
213 combined approaches.

214

215

Correlative approaches

Distributional changes are typically estimated through the use of correlative models that aim to represent the realized niche of a species[48,49]. Correlative models relate observed geographic distribution of a species to current climate; resultant models are then applied to climate projections to infer potential climatically-suitable areas for a given species in the future. Species' distribution can be presence-only data[17,22], presence/absence[50] or abundance observations[51], based either on fieldwork or specimen records[22,52]. Correlative models have been applied to species at scales ranging from local to global[19,53] (Fig. 1), and have been widely used to explore the vulnerability of vertebrates (including birds[36,52,54], mammals[17,28], amphibians[30,50], fishes[22,55]), invertebrates[14,56,57] and plants[27,58].

Correlative models have the advantage of being spatially explicit and they are applicable to a wide range of taxa at various spatial scales. However, there are a number of limitations and uncertainties associated with them (see Pearson *et al.*[29] and Wiens *et al.*[59] for detailed descriptions). Primary sources of uncertainty and potential errors can be divided into three broad classes: climatic, algorithmic, and biotic[29,59]. Climatic uncertainties, that apply to all types of approaches, may arise from general circulation models, which use different parameters and model structures to simulate future climate systems, and may produce different results irrespective of the assumed greenhouse gas emissions[59,60]. Climate models project future climate conditions at a coarser scale of resolution than that of data (biological and environmental) used to calibrate the correlative models[49,59], and their outputs are thus often not sufficiently fine-scaled for modelling rare species or species with small geographic distributions[49,50]. Algorithmic uncertainties can arise from the differences in methods and models used to predict species' distribution (e.g. Generalized Additive Models, Maximum Entropy models), and from the selection of model predictors (e.g. mean annual temperature, annual precipitation; see[61]), which have shown great variability in both results and model performance. This range of uncertainties has been addressed by some by applying a variety

241 of different statistical methods and model structures, summarising predictions across all models to
242 generate ensemble forecasts, e.g. model-averaged probability of presence and confidence intervals
243 (see examples[16,30,62]). Biotic uncertainties may arise if the assumptions made about a species'
244 biology are inappropriate. First, species' distributions are assumed to be in equilibrium with
245 surrounding climates and these relationships are assumed to persist in the future[56]. Second, it is
246 unknown how much of a species' fundamental niche, exclusively determined by the species'
247 requirements and/or tolerances is represented by its currently realized niche, which is also
248 determined by abiotic, biotic, geographic, historical and anthropogenic factors[49]. Moreover,
249 correlative models for plants do not account for drivers such as changes in atmospheric CO_2
250 concentration, which influence plant growth and water use and can alter demographic processes
251 sufficiently to drive ecosystem structural and functional changes[63]. Correlative models can also be
252 used to predict future geographic distribution of a group of species in a given area and the results
253 combined to create assessments of new community structures[64]. However, these models ignore
254 community-assembly rules, as well as differences in the constraints and adaptability of individual
255 species, and thus the resulting predicted species assemblages may be unrealistic[62]. Correlative
256 models have been criticised by some authors because they lack mechanism and causality (e.g.
257 see[65]), although there is increasing evidence that recent population trends have matched those
258 expected from correlative model projections[36].

259 The relatively large number of reliable occurrence points required to fit correlative models
260 often precludes their use for assessments of poorly known species[66]. They are also less appropriate
261 for species with cosmopolitan or limited geographic distributions (e.g. on small islands) since
262 climate may not explain distributions or distributional changes. Despite these limitations, the
263 majority of regional and global analyses to date are based on correlative approaches, since they can
264 be relatively quick and cheap to apply[67] and occurrence data are available for a large number of
265 taxa.

Mechanistic approaches

Mechanistic models require taxon-specific parameters that provide information on the behaviour of individuals and the mechanisms they develop to cope with changing climatic conditions. Mechanistic models are developed from laboratory and field observations of demographic rates, physiological tolerances[41,68,69], competition and dispersal[70], diseases and predation[71], as well as from energy balance equations[72]. Measures of vulnerability derived from these models are typically expressed in terms of probability of extinction, whether of discrete populations or entire species. Mechanistic approaches often focus on a single species of conservation interest (e.g. rare or threatened species)[39,41], since methods used to collect detailed data on species physiology, which are essential to parameterise such models, are costly and time-consuming. Some studies exist involving this type of modelling that do not involve a specific taxon but rather provide general theoretical frameworks to predict effects of climate change on plants[10], terrestrial ectotherms[68] and generic species[9,10], highlighting major determinants of extinction risk in a changing environment and providing recommendations for future research needs. Some mechanistic models (e.g. incidence function models, age-structured metapopulation models) may be used to explain metapopulation dynamics in the presence of climate change by estimating extinction and colonization rates as functions of habitat suitability[73], prey availability or management actions[45]. Other mechanistic models consider the changes in vegetation distribution and dynamics using both bioclimatic and physiological parameters of groups of species (e.g. plant functional types) [74].

Mechanistic niche models utilise species' functional traits, physiological tolerances and energy and mass exchanges to represent the fundamental niche of a species[75]. Key functional traits (e.g. morphology, physiology, behaviour) and spatial habitat data (e.g. climate, vegetation cover, topography, bathymetry) are used to assess individual fitness[75,76]. Such models are considered by some authors to be more robust and theoretically defensible than correlative models for predicting

291 species' responses to climate change[75]. Compared to the realized niche modelled via correlative
292 approaches, the mechanistically modelled fundamental niche provides a better approximation of the
293 climatic space in which an organism can exist, including areas that have, or may, become newly
294 suitable[75,76]. In addition, these models permit explicit consideration of important biological factors
295 like evolutionary changes and physiological responses.

296 Extensive application of mechanistic niche models is precluded by the fact that they require
297 detailed data that are lacking for most species. The main sources of uncertainty in mechanistic
298 models relate to model parameters (e.g. population abundance, which may be underestimated
299 depending on the method used to collect the data and the ability of the observer to detect the
300 species), and to combining data collected at different spatial resolutions[23]. Moreover, these models
301 usually do not account for non-climatic threats to dispersal or for biotic interactions[48].

302

303 **Trait-based vulnerability assessment approaches**

304 TVAs use species' biological characteristics as predictors of extinction risk due to climate
305 change[13,14], often in combination with estimates of exposure. Methods typically involve selecting
306 traits related to sensitivity (e.g. typically describing ecological specialization, inter-specific
307 interactions) and adaptability (i.e. dispersal and phenotypic adaptability[14,77,78]) and scoring each
308 according to observations or expert judgment [79,80]. For example, Gardali *et al.*[78] quantified the
309 vulnerability of Californian birds by scoring sensitivity and exposure for each taxon. They used
310 information from published literature to assign a sensitivity score to four intrinsic species'
311 characteristics (dispersal ability, migratory status, habitat specialization and physiological
312 tolerances), and then combined sensitivity and exposure scores to generate a climate vulnerability
313 index.

TVAs are being used increasingly by conservation organizations and management agencies because they permit a relatively rapid assessment for multiple species, which can be used to prioritize conservation planning and implementation of adaptation schemes. Moreover, TVAs are sometimes considered easier to use by practitioners because they do not require extensive knowledge of modelling techniques, even if their applicability is limited to a specific area and to cases where relevant data on species' traits are available (see[81]).

Drawbacks with TVAs are that precise vulnerability thresholds associated with each trait are often unknown, necessitating selection of arbitrary, relative thresholds for categories of higher or lower extinction risk. Traits are often weighted equally[20] even though some characteristics are likely to be more important than others in determining climate change vulnerability. Subject to the challenges of score-based systems, it is not possible to compare vulnerability between taxonomic groups for which different sets of traits may have been used in the TVA. Moreover, different TVAs applied to the same species do not always yield congruent results[82]. The most common sources of uncertainty in TVAs stem from the choice of traits included in assessments, parameterisation of thresholds of associated vulnerability, and from gaps in knowledge of individual species' characteristics[14,83]. For example, dispersal distance is one of TVA's most important and conservation-informative traits, yet estimates are currently available for few animal species. Some studies have attempted to provide dispersal estimates[16,18,84], but inevitable uncertainties arise from models and parameters. Uncertainty is usually incorporated as a confidence score based on expert opinion. Such score can be provided for each trait[78], for each stage of the assessment[83], or for the overall assessment[78]. Alternatively, some authors rank missing trait data under best- or worst-case scenarios[14,80], by assuming optimistic and pessimistic extreme values.

337 **Combined approaches**

338 There is a growing consensus on the benefits of using approaches that combine different
339 types of models and data[32,40]. Here we discuss the three most common combined approaches,
340 criteria-based, mechanistic-correlative and correlative-TVA.

341

342 **Criteria-based approaches**

343 Criteria-based approaches have been used to combine observed or projected demographic
344 trends (e.g. population increases or decreases) with intrinsic characteristics of species (e.g.
345 generation length), to classify species into threat categories based on the risks posed by climate
346 change. Climate-attributed changes in species' geographic ranges, often derived from correlative
347 models, are assessed against quantitative thresholds[34,38,83,85]. These assessments often use the IUCN
348 Red List categories and criteria (www.iucnredlist.org)[38,85] or draw inspiration from them[83].

349 One advantage of criteria-based approaches is that they can be applied to large numbers of
350 species worldwide[86]. They are important for assessing the conservation status of species threatened
351 by climate change since they simultaneously account for several factors known to affect the relative
352 extinction risk (e.g. declines in the extent of occurrence, reduction in population size). Furthermore,
353 by using quantitative thresholds to predict relative extinction risk, it is possible to make
354 comparisons between past, current and future conservation status of species. Approaches based on
355 the IUCN Red List require a consistent adoption of thresholds and criteria[87]; however, these are
356 sometimes arbitrarily modified (e.g. to temporal and spatial scales and spatial resolution), thereby
357 reducing the comparability and interpretability of the results[87]. Pearson *et al.*[88] identified factors that
358 predispose a selection of North American herpetiles to high extinction risk due to climate change,
359 and concluded that most important factors are already incorporated into extinction risk assessments
360 for the IUCN Red List.

Mechanistic-correlative and mechanistic-correlative-TVA approaches.

In mechanistic-correlative approaches, outputs of correlative models are incorporated into demographic models to calculate spatial structure of populations[45], whose dynamics are then modelled mechanistically. This combination is useful, for example, in predicting how distribution patterns influence the viability of populations under a changing climate[32,40]. Furthermore, some studies have integrated life-history characteristics into models to produce more accurate projections of species' responses to climate change. Keith *et al.*[32] assessed extinction risk for plant species in South African fynbos under stable and changing climatic conditions. The authors linked the outputs of correlative models with a demographic metapopulation model, and considered their interactions with fire tolerances and dispersal abilities. In this way, they dealt with both habitat changes and population dynamics simultaneously in their assessments.

Correlative-TVA approaches

Other combined approaches integrate species characteristics and species distribution models by incorporating species traits to refine distribution projections made using correlative models[16,18,31,89], or by integrating correlative model outputs into trait-based assessments[21,83]. In the first approach, traits like dispersal ability and generation length have been usefully applied to refine range dynamics[16,90]. For example, Barbet-Massin *et al.*[16] used natal dispersal and generation length to predict the breeding distribution of European birds under climate and land-use changes. The authors predicted a 10% reduction of future species richness assuming unlimited dispersal and a 25% reduction by using natal dispersal.

In the second type of approach, the outputs of correlative models are used to estimate exposure to climate change and identify areas, which might become suitable in the future, even if they fall outside a species' current range. By linking exposure, estimated with correlative models,

385 with sensitivity and adaptability assessed with TVAs, a vulnerability index can be calculated that
386 accounts for both intrinsic and extrinsic factors (e.g.[83]).

387

388 **Guidance for selecting climate change vulnerability assessment approaches**

389 Ideally, practitioners should assess the vulnerability of populations or species to climate
390 change using a variety of methods, with greatest predictive confidence conferred where models are
391 in agreement. The choice of the approach is entirely dependent on conservation goals, which are
392 often vague and not clearly defined, and on the data available (Box 1). Relying on these broad
393 goals, practitioners need to identify definable and measurable objectives[91], in terms of temporal,
394 spatial and taxonomic scales. In Table 1 we identify different examples of objectives against each
395 approach and below provide two exemplary goals and identify the associated methodologies to
396 reach them.

397

398 **Estimating extinction risk**

399 When deriving estimates of extinction risk of species is the goal, both mechanistic and
400 correlative models can provide appropriate results. The most effective way to predict extinction risk
401 of species under climate change is to combine demographic data (e.g. population trends, survival,
402 fertility) with changing environmental factors (e.g. precipitation, sea ice extent), and then project
403 these changes into the future[41,43]. For example, Jenouvrier *et al.*[41] used a mechanistic model, which
404 combined demographic and climatic data, to project a > 35% probability of extinction for the
405 emperor penguins (*Aptenodytes forsteri*) in Antarctica by 2100 in response to projected sea ice
406 changes.

407 Another way of inferring the extinction risk of species is to use a decline in suitable area as a
408 proxy for population decline[38,92,93], providing the relationship between the two can be assumed to
409 remain constant. Correlative models can be used to project range changes into the future; this would

allow classifying the species into one of IUCN Red List categories. Levinsky *et al.*[93], for example, demonstrated that the proportion of European mammals that are forecast to become extinct by 2100 can vary from 1 to 9%, depending on the magnitude of predicted climatic changes and the ability of species to migrate.

Prioritization of actions

Climate change adaptation strategies require creating a link between an explicitly stated expectation about the way global warming could affect species, habitats, or even people, to clear objectives and actions that would best address those climate impacts[94]. Conservation decision-making is about prioritizing actions to satisfy conservation objectives for a set of species and areas[95]. It is not possible to make conservation interventions for all species, and prioritization exercises are needed to determine which actions to focus on to protect species[96]. Given the high levels of uncertainty and complexity in modelling impacts, we highlight that reprioritizing or even abandoning actions which benefit some species over others should be done with great caution.

Where site-scale conservation is the focus (e.g. in a protected area), correlative models are able to identify species for which the area may be suitable in the future, thereby allowing managers to prepare for potentially novel species assemblages and plan appropriate conservation actions (e.g. predator and invasive species control). For example, Hole *et al.*[54] used correlative models to assess species turnover in a network of Important Bird Areas in Africa, and provided generic guidance on the types of conservation actions (e.g. translocation, habitat restoration, disturbance-regime management) that might be most appropriate for individual sites.

For a regional-scale focus, identifying the bioclimatic space where species could persist and the areas of relatively unchanged climate within this space may facilitate species persistence during periods of climatic stress. Spatially explicit projections from correlative and mechanistic niche models allow identification of these sites. For example, Maschinski *et al.*[43] used a mechanistic

approach to identify potential climatic refugia for an endemic plant species (*Purshia subintegra*) of Arizona. This study showed that in situ manipulation and introductions at northern latitudes are priority actions necessary to prevent the extinction of this rare and endangered species.

Where the focus is on particular species, trait-based and mechanistic approaches are likely to deliver insights into the specific mechanism(s) of impact (e.g. increased competition, loss of mutualisms, disruption of cues, disease)[14], allowing targeted interventions both to decrease species' sensitivity (e.g. disease treatment, predator control) and to increase their adaptive capacity (e.g. genetic management, improved landscape permeability, translocation)[75]. Indices calculated with trait-based approaches can facilitate grouping taxa by their relative risk to climatic changes, which help identify adaptation strategies that could benefit multiple species[77]. For example, Moyle *et. al*[80], who assessed Californian freshwater fishes according to their life-history characteristics, classified species that were heavily dependent on human intervention as highly vulnerable to climate change, and highlighted the need for conservation actions such as management of barriers, special flows and removal of alien species to allow population persistence.

Conclusions

This review of climate change vulnerability assessment approaches suggests that, in general, a correlative approach is appropriate when the only data available are those on species' occurrence, in particular for reconstructing the paleoclimatic niche of fossil species or projecting their future climatic suitable area, from local to global scales. On the other hand, mechanistic models have the greatest power to assess extinction probability driven by climate change, identify conservation actions and evaluate the potential effectiveness of management interventions, but they are limited to few terrestrial species. Therefore, they are usually employed when the focus is on a well-studied species of particular conservation interest (e.g. species threatened, keystone, flagship or umbrella), for which detailed physiologic and/or demographic data are available. Trait-based approaches are

460 less resource-intensive and therefore more widely used. This method is ideal to help non-GIS
461 experts develop regional assessments and to identify conservation priorities in the absence of
462 specific data on species' distribution.

463 Validation of the accuracy and precision of vulnerability assessment approaches, through
464 comparison of model projections with a globally coordinated observation effort, is essential for
465 improving projections of the impacts of climate change on species. Use of paleoecological evidence
466 of past species' responses to climatic variation in conjunction with matching paleoclimatic data can
467 provide an opportunity to test the assessments[97,98]. Observations of recent responses to climate
468 change are another useful tool to test reliability of model predictions against current observations.
469 However, quantifying the ability of models to provide reliable range shift projections or population
470 changes is still challenging, since they are often difficult to validate across time and space[97]. One
471 key issue is the debate on modelling the realized vs. the fundamental niche[48,49,79]. Both the lack of
472 equilibrium between species and climate, and the difficulty of isolating the effects of climatic
473 changes on a species' range from those of other threats[97], can lead to changes in the realized niche
474 of a species (usually modelled mechanistically). On the other hand, correlative approaches attempt
475 to model the fundamental niche of a species, but they use data from the realized niche[48]. This can
476 lead to spurious correlations between species' occurrence and climate and thus hinder model
477 validation as well as casting doubts on model accuracy[48]. For example, a species may not respond to
478 climate only because other factors (e.g. competitive exclusion, predation) are confounding the
479 response[99]. Additionally, when comparing past and current distribution to validate models or TVAs,
480 a big challenge is to find accurate information on species' historic distribution and population
481 trends. Addressing all of these issues should lead to better conservation decision-making.

482 A glaring oversight in almost all studies is that they only focused on the direct impacts of
483 climate change. Indirect impacts within biological communities, as well as changes in human use of
484 natural resources are going to have substantial, complex, and often multiplicative impacts on

485 species[36,100]. Thus, many current assessments are blind to the fact that the interactions between
486 current threats and climate change are likely to be profound[3]. Moreover, the growing human
487 population will itself be increasingly affected by climate change, with human adaptation responses
488 likely to result in substantial and negative impacts on biodiversity[100]. Assessments of future impacts
489 of climate change need to take these factors into account.

490

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723

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727

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729 M.P., P.V., C.R., J.E.M.W. and W.B.F. designed the framework for the review. All authors contributed to the
730 writing, discussed the results and commented on the manuscript.

731

732 **Competing financial interests**

733 The authors declare no competing financial interests.

734

735 **Figure headings:**

736

737 **Figure 1:** Taxonomic focus of vulnerability assessments in the analysed papers.

738 Birds are the most analysed taxon, followed by mammals and plants, while invertebrates other than
739 insects have seldom been assessed. Colours represent the spatial scale of the assessments. Regional
740 scale is defined as describing the range of 10^4 to 10^7 km^2, while scales smaller than 10^4 km^2 are
741 referred to as local scales.

742 **Figure 2:** Ecoregional global concentrations of terrestrial and marine climate change vulnerable
743 species.

744 Studies conducted at regional, continental and global scales where used to derive a global map of
745 vulnerability, according to an ecoregional classification. The red scale represents terrestrial areas
746 with high numbers of vulnerable species, identified on the basis of 1) the number of species
747 assessed and 2) the taxonomic ranks higher than species considered. The blue scale represents areas
748 that host marine vulnerable species. Dark colours indicate areas of high vulnerability, while light
749 colours indicate areas of relatively low vulnerability.

750 **Figure 3:** Trends and biases in taxonomic groups assessed and approaches used by continent.

751 Birds and mammals have been the most frequently analysed taxa across all continents between 1995
752 and 2014, usually with similar proportions (with the exception of Asia). Correlative approaches are
753 widely used for assessing species vulnerability in Africa, Asia and Europe, while mechanistic
754 approaches prevail in North America. Trait-based approaches are used mostly in Australia and
755 North America.

756

757 **Table 1 | Examples of objectives in climate change vulnerability assessments, on the basis of the scale to be adopted.**

	Temporal scale			Spatial scale			Taxonomic scale		
	Past	**Recent past/ present**	**Present/ Future**	**Local/site**	**Regional**	**Global**	**Population and ranks < than species**	**Single species**	**Multispecies**
Examples of objectives: correlative	Reconstructing species' past distribution[101]	Modelling current climatic suitable areas for species[22]	Predicting climate-induced future range shifts under different time intervals[102]	Quantifying the area that will remain climatically suitable for species living in areas important for conservation[60]	Assessing the ability of a network of protected areas to ensure the persistence of species[103]	Identifying the most important climatic variables in determining a species' distribution globally[19]	Quantify the latitudinal/ altitudinal shifts of the various populations of a species[104]	Assessing a species' future threat status[93]	Predicting spatial patterns of species richness[105]
	Identifying past climatic refugia[106]	Quantifying % range gains/losses in the last decades to estimate extinction risk[38]	Projecting future range margin contractions/ expansions by 2080[92]	Quantifying species' turnover within a protected area[54]	Identifying and designing potential areas to be protected within a region[107]	Identifying hotspots of species highly exposed[19]	Assessing which of the populations of a species will experience the greatest changes in its distribution[104]	Predicting spatial overlap between the current and future range of a species[108]	Modelling possible future community assemblages[109]
Examples of objectives: mechanistic	Representing postglacial expansions from glacial refugia[110]	Quantifying population reductions in recent times due to changes in sea ice extent[41]	Predicting survival under future climate change[111]	Determining climatic factors that affect reproductive success of a reintroduced species[112]	Exploring the range margin dynamics for species of conservation concern within a region[40]	Assessing species thermal tolerances across their range[113]	Assessing the extinction risk of a population at the margins of a species' range[40]	Assessing the impacts of sea level rise on a coastal species[114]	Modelling prey-predator dynamics under future climatic conditions[45]

	Understanding the effects of changes in CO_2 concentration on plants[115]	Determining population viability due to an increase in frequency of extreme climatic events during the last decades[43]	Assessing species' probability of extinction by 2100[41]	Predicting the probability of extinction of a keystone species within a site[42]	Exploring the extinction risk of a species in part of its range[39]	Predicting changes in fitness due to global warming globally[68]	Determining the extinction risk of a threatened subspecies[34]	Estimating species' abundance in the future under climate change[116]	Predicting future expansions of invasive species[117]
Examples of objectives: TVA	Identifying trends of past extinctions related to life history traits[118]	Identifying taxonomic groups that currently retain high numbers of sensitive and unadaptable species[78]	Identifying sensitive species living in areas that are likely to become highly exposed in the future[119]	Prioritizing conservation actions at the local scale[120]	Making an assessment of species vulnerability within a country[80]	Identifying species with the greatest relative vulnerability to climate change[78]	Identifying potential adaptive characteristics of an isolated subspecies[35]	Identifying the traits that make a species most vulnerable to climate change[120]	Identifying the most vulnerable species to climate change within a taxon[20]
	Predicting the response of species, that share life history traits with past extinct/impacted species, to future climatic changes[121]	Identifying the characteristics of species that played the most important role in determining reductions/ extinctions in recent years[14]	Identifying unadaptable species with the largest predicted range shifts in the coming decades[83]	Understanding which component of vulnerability is prevalent for a species within a site[122]	Understanding how traits relate to changes in occurrence of species within a freshwater basin subject to droughts[11]	Identifying areas with the greatest number of vulnerable species at the global scale[14]	Identifying potentially vulnerable subspecies/ populations/ varieties with relatively unknown distribution[36]	Assessing species' adaptive capacity/ resilience[14]	Selecting different adaptation strategies according to the relative vulnerability of different species[78]

*References from 101 to 122 are listed in the Supplementary material.

Box 1 | Data availability

Once clear objectives have been established, and the potential approaches identified, another consideration for selecting the most appropriate method is to consider the types of data available. The financial resources, time, expertise and input data required for each method are likely to mean that just one or, at best, a few approaches are feasible. When fine scale data on species occurrence are available (e.g. point localities), correlative and mechanistic niche models may be applied. To build these types of models, adequate climate data covering different time periods are also needed. For example, paleoclimatic reconstructions for Paleocene and Holocene, as well as current and future projections, are already available under different resolutions and time intervals (e.g.[123,124]).

Where relevant life-history data (e.g. data on species' biology, ecology, physiology, demography) are available; (see ecology and trait databases for birds[125], mammals[81,126] and amphibians[127]) , trait-based or mechanistic approaches could facilitate, for example, the identification of resilient and/or adaptable species, thus aiding in prioritization[11]. Moreover, these kinds of data are necessary to develop mechanistic niche models to refine species' distribution based on the mechanisms that species themselves develop to cope with global warming[13]. Often this type of empirical data will be lacking. Rather than abandon modelling and informing conservation decisions in these cases, structured expert elicitation approaches offer an interim way of estimating key species demographic and life-history parameters[128,129] .